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# A TOEPLITZ GRAM-SCHMIDT ALGORITHM FOR AUTOREGRESSIVE MODELING

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#### INTRODUCTION

Let  $\{y(t), t \in Z\}$  be a zero mean, covariance stationary, discrete time series with autocovariance function  $\{R(v), v \in Z\}$ . Then in: 1) finite order autoregressive modeling (Akaike (1969)), 2) spectral density estimation using autoregressive approximants (Parzen (1974)), 3) maximum entropy spectral analysis (Ulrych and Bishop (1974)), and 4) explicit minimum mean squared error linear prediction (see Anderson (1971), p. 419), one is faced with estimating the quantities  $\alpha_k(j), \sigma_k^2, 1 \le j \le k \le M$ , satisfying the Yule-Walker equations

$$\sum_{j=0}^{k} \alpha_{k}(j) R(j-v) = \delta_{v} \sigma_{k}^{2} , v \ge 0 , \qquad (1.1)$$

where  $\alpha_k(0) = 1$  and  $\delta_v$  is the Kronecker delta.

Let y(1),...,y(T) be a mean detrended sample realization from y. The traditional method for finding estimators  $\hat{\alpha}_k(j)$  and  $\hat{\sigma}_k^2$  of  $\alpha_k(j)$  and  $\sigma_k^2$  has been to estimate R(0),...,R(M) by the positive definite sample autocovariances

$$\hat{R}(v) = \frac{1}{T} \sum_{t=1}^{T-|v|} y(t)y(t+|v|) , |v| \leq M,$$

and then to use (1.1) with  $\hat{R}(v)$  replacing R(v). These Yule-Walker estimators have two main attractions; 1) they can be calculated very quickly by Levinson's (1947) algorithm

$$\hat{\alpha}_{1}(1) = -\hat{R}(1)/\hat{R}(0)$$
 ,  $\hat{\sigma}_{1}^{2} = \hat{R}(0)[1-\hat{\alpha}_{1}^{2}(1)]$  (1.2)

$$\hat{\alpha}_{k}(k) = -\left[\sum_{j=0}^{k-1} \hat{\alpha}_{k-1}(j)\hat{R}(k-j)\right]/\hat{\sigma}_{k-1}^{2} ,$$

$$\hat{\alpha}_{k}(j) = \hat{\alpha}_{k-1}(j) + \hat{\alpha}_{k}(k)\hat{\alpha}_{k-1}(k-j), \qquad j=1,...,k-1,$$

$$\hat{\sigma}_{k}^{2} = \hat{\sigma}_{k-1}^{2} \left[1-\hat{\alpha}_{k}^{2}(k)\right] ,$$
(1.3)

k=2,...,M, and 2) they are guaranteed to lead to stable prediction filters and positive spectral density estimators since (Parzen (1961)) the zeros of the complex polynomials  $\hat{g}_k(z) = \sum_{i=0}^k \hat{\alpha}_k(j)z^i$  are all outside the unit circle.

If we define the  $(k \times k)$  Toeplitz matrix  $\hat{\Gamma}_k$  to have (i,j)th element  $\hat{R}(|i-j|)$  and the  $(k \times 1)$  vector  $\hat{\Gamma}_k$  to have  $i^{th}$  element  $\hat{R}(i)$ , we can write the sample Yule-Walker equations for  $v=1,\ldots,k$  as

$$\hat{\Gamma}_{\mathbf{k}} \hat{\alpha}_{\mathbf{k}} = -\hat{\Gamma}_{\mathbf{k}} \qquad , \tag{1.4}$$

where  $\hat{q}_k = (\hat{q}_k(1), \dots, \hat{q}_k(k))^T$ . Then (1.4) are the normal equations for the ordinary least squares estimators  $\hat{q}_k$  in the linear model  $y_k = -X_k x_k + \varepsilon$ , where

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$$y_{k} = \begin{bmatrix} y(1) \\ y(2) \\ y(3) \\ \vdots \\ y(k+1) \\ \vdots \\ y(T) \\ 0 \\ 0 \\ \vdots \\ 0 \end{bmatrix} , X_{k} = \begin{bmatrix} 0 & 0 & \dots & 0 \\ y(1) & 0 & \dots & 0 \\ y(2) & y(1) & \dots & 0 \\ \vdots \\ y(k) & y(k-1) & \dots & y(1) \\ \vdots \\ y(T) & y(T-1) & y(T-2) & \dots & y(T-k) \\ y(T) & y(T-1) & \dots & y(T-k+1) \\ 0 & y(T) & y(T-1) & \dots & y(T-k+2) \\ \vdots & \vdots & \vdots & \vdots \\ \vdots & \vdots & \vdots & \vdots \\ 0 & 0 & \dots & y(T) \end{bmatrix}$$
 (1.5)

are (T+k)  $x^-1$  and (T+k) x k respectively. Thus the Yule-Walker estimators correspond to the ordinary least squares regression of y(t) on  $y(t-1), \ldots, y(t-k)$  for  $t=1,\ldots,T+k$ ; with any unobserved y replaced by its expected value of zero.

Thus the Yule-Walker estimators suffer numerically by being the solution to normal equations for a linear model rather than being the result of applying a triangularization method (see Lawson and Hanson (1974), for example) to the design matrix  $X_k$ . This can be particularly important in time series because of ill-conditioning caused by the peculiar form of  $X_k$ . In addition, if the variance of y is large, then the effect of replacing the observations  $y(1-k), \ldots, y(0)$  and  $y(T+1), \ldots, y(T+k)$  that are on the ends of the observed data by zeros raises the possibility of obtaining poor estimators.

There have been three main attempts to alleviate the problems in Yule-Walker estimators. First, are what are often called least squares estimators (see Anderson (1971), p. 211). These are obtained by dropping the first and last k rows of  $y_k$  and  $X_k$  and then performing ordinary least squares regression.

These estimators seem to reduce end effects but cannot be obtained recursively for  $k=1,\ldots,M$  and also are not guaranteed to lead to stable filters. The second method is the Burg (1968) algorithm which seems to moderate end effects, is recursive in nature, and leads to stable filters. However, it appears to be somewhat ad hoc in nature and its numerical properties are difficult to describe. The third method, due to Pagano (1972), consists of solving the normal equations (1.4) via the numerically stable modified Cholesky triangularization (Wilkinson (1967)) of  $\hat{\Gamma}_M$ .

The purpose of this paper is to show how the factors needed to find the modified Gram-Schmidt triangularization of  $X_M$  corresponding to Pagano's triangularization of  $\hat{\Gamma}_M$  can be found recursively while only needing to store two (T+M) x l vectors (rather than storing the entire  $X_M$  matrix). From this we find a design-matrix oriented algorithm, called the Toeplitz Gram-Schmidt algorithm, for finding the Yule-Walker estimators. We show that the Toeplitz Gram-Schmidt algorithm is roughly as fast as using Levinson's algorithm, while providing a unified framework for deriving other stable, more robust estimators. The algorithm allows calculation of diagnostics useful in measuring end effects as well as determining if particular observed values of y are exerting undue influence on the estimation procedure.

The algorithm is given in section 2, its numerical properties described in section 3, and its application for robust autoregression and autoregression diagnostics discussed in section 4.

## 2. THE TOEPLITZ GRAM-SCHMIDT YULE-WALKER ALGORITHM

We seek ordinary least squares estimators of  $\alpha_k$ , k=1,...,M, for the linear models  $y_k = -X_k \alpha_k + \varepsilon$  without using the normal equations.

By Levinson's algorithm we can obtain all the  $\hat{\alpha}_k(j)$  and  $\hat{\sigma}_k^2$ , from only  $\hat{R}(0)$  and  $\hat{\alpha}_k(k)$ ,  $k=1,\ldots,M$ , by

$$\hat{\sigma}_1^2 = \hat{R}(0) \left[1 - \hat{\alpha}_1^2(1)\right] \tag{2.1}$$

$$\hat{\alpha}_{k}(j) = \hat{\alpha}_{k-1}(j) + \hat{\alpha}_{k}(k)\hat{\alpha}_{k-1}(k-j)$$
,  $j=1,...,k-1$ , (2.2)

$$\hat{\sigma}_{k}^{2} = \hat{\sigma}_{k-1}^{2} \left[1 - \hat{\alpha}_{k}^{2}(k)\right], \tag{2.3}$$

k=2,...,M. Further if one only wants to estimate  $\sigma_k^2/R(0)$ , as is often the case, then only  $\hat{\alpha}_k(k)$ , k=1,...,M are required.

Thus the algorithm we propose consists of determining recursively  $\hat{\alpha}_k(k)$ ,  $k=1,\ldots,M$ , by calculating at the kth step the kth and (M+1)st columns of the kth stage of the modified Gram-Schmidt decomposition (Björck (1967)), MGSD, of the  $(T+M) \times (M+1)$  matrix  $[X_M; y_M]$ . Then these  $\hat{\alpha}_k(k)$ 's are used in (2.1)-(2.3) to find all the desired estimators. Theorem 2.1 shows how the  $\hat{\alpha}_k(k)$ 's are calculated while Theorem 3.1 shows that the algorithm is indeed an application of the MGSD.

Define the lag operator L on the  $(n \times 1)$  vector  $\mathbf{a} = (\mathbf{a}(1), \dots, \mathbf{a}(n))^T$  to be  $L(\mathbf{a}) = (0, \mathbf{a}(1), \dots, \mathbf{a}(n-1))^T$ . Then we note that the columns of  $X_M$  are  $x_1, \dots, x_M$  where  $x_1 = L(y_M)$  and  $x_j = L(x_{j-1})$ ,  $j=2,\dots,M$ .

## Theorem 2.1

Let 
$$y_{(1)} = y_M$$
,  $x_{(1)} = L(y_M) = x_1$ , and for  $k = 1, ..., M$ :

$$\hat{a}(k) = -\frac{y \left(k\right) \times (k)}{T}$$

$$\times (k) \times (k)$$

$$\frac{y}{z}(k+1) = \frac{y}{z}(k) + \hat{a}(k) \times (k)$$

$$x(k+1) = L(x(k) + \hat{a}(k)y(k))$$
.

Then

a) 
$$\hat{a}(k) = \hat{\alpha}_{k}(k)$$
 ,  $k=1,...,M$ .

b) 
$$y_{(k)} = y_M + \sum_{j=1}^{k-1} \hat{\alpha}_{k-1}(j) x_j$$
,  $k=1,...,M+1$ ,

c) 
$$x_{(k)} = x_k + \sum_{j=1}^{k-1} \hat{\alpha}_{k-1}(j) x_{k-j}, k=1,...,M+1,$$

where any summation is zero if its upper limit is less than its lower limit.

# Proof

We proceed by induction on k. For k=1, (b) and (c) are true by definition, while  $\hat{a}(1) = -T\hat{R}(1)/T\hat{R}(0) = \hat{\alpha}_1(1)$ . For k>1, we have by the induction hypothesis,

$$x(k) = L[x(k-1) + \hat{a}(k-1) y(k-1)]$$

$$= L[x_{(k-1)}^{k-2} + \sum_{j=1}^{k-2} \hat{\alpha}_{k-2}^{j} + \sum_{j=1}^{k-2} \hat{\alpha}_{k-1}^{j} + \hat{\alpha}_{k-1}^{k-1} + \sum_{j=1}^{k-2} \hat{\alpha}_{k-2}^{j} + \hat{\alpha}_{k-1}^{j} + \hat{$$

by (1.3). Similarly,

$$\begin{aligned} y_{(k)} &= [y_{M} + \sum_{j=1}^{k-2} \hat{\alpha}_{k-2}(j) x_{j}] + \hat{\alpha}_{k-1}(k-1)[x_{k-1} + \sum_{j=1}^{k-2} \hat{\alpha}_{k-2}(j) x_{k-1-j}] \\ &= y_{M} + \hat{\alpha}_{k-1}(k-1) x_{k-1} + \sum_{j=1}^{k-2} [\hat{\alpha}_{k-2}(j) + \hat{\alpha}_{k-1}(k-1) \hat{\alpha}_{k-2}(k-1-j)] x_{j} \\ &= y_{M} + \sum_{j=1}^{k-1} \hat{\alpha}_{k-1}(j) x_{j} \end{aligned} .$$

We have thus shown (a)-(c) being true for k-1 implies (b) and (c) are true for k and it remains to prove (a). To do so we need several well-known facts about the modified Cholesky decomposition of  $\hat{\Gamma}_{M}$  and the Gram-Schmidt decomposition of  $X_{M}$ .

Let  $X_M = Q_M R_M$  be the Gram-Schmidt decomposition of  $X_M$ , i.e  $R_M$  is an  $(M \times M)$  unit upper triangular matrix and  $Q_M$  is a  $(T + M) \times M$  orthogonal matrix having columns  $q_1, \dots, q_M$ , i.e  $Q_M^T Q_M = D_M$ , a diagonal matrix. Let  $\hat{\Gamma}_M = \hat{L}_M \hat{D}_M \hat{L}_M^T$  be the modified Cholesky decomposition of  $\hat{\Gamma}_M$ , i.e  $\hat{L}_M$  is an

(M x M) unit lower triangular matrix and  $\hat{D}_{M}$  is am (M x M) diagonal matrix. Then  $\hat{\Gamma}_{M} = \frac{1}{T} X_{M}^{T} X_{M} = \hat{L}_{M} \hat{D}_{M} \hat{L}_{M}^{T} = \frac{1}{T} R_{M}^{T} D_{M} R_{M}$ , which by the uniqueness of the modified Cholesky decomposition (Wilkinson (1967)), gives  $\hat{L}_{M} = R_{M}^{T}$  and  $\hat{D}_{M} = \frac{1}{T} D_{M}$ . Further (Pagano (1972)), with  $\hat{\sigma}_{O}^{2} = \hat{R}(0)$ , we have

$$\hat{L}_{M}^{-1} = \begin{bmatrix} 1 & & & & \\ \hat{\alpha}_{1}(1) & 1 & & & \\ \hat{\alpha}_{2}(2) & \hat{\alpha}_{2}(1) & 1 & & & \\ \vdots & & \ddots & & \vdots \\ \hat{\alpha}_{M-1}(M-1)\hat{\alpha}_{M-1}(M-2)\dots\hat{\alpha}_{M-1}(1) & 1 \end{bmatrix} , \qquad (2.4)$$

$$\hat{D}_{M} = Diag \left(\hat{\alpha}_{0}^{2}, \hat{\sigma}_{1}^{2}, \dots, \hat{\sigma}_{M-1}^{2}\right) ,$$

and, defining  $\hat{\theta}_{M} = (\hat{\alpha}_{1}(1), \dots, \hat{\alpha}_{M}(M))^{T}$ ,

$$\hat{\theta}_{M} = -(Q_{M}^{T}Q_{M})^{-1} Q_{M}^{T} Y_{M} \qquad ,$$

<u>i . e</u>

$$\hat{\alpha}_{k}(k) = -\frac{g_{k}^{T} Y_{M}}{g_{k}^{T} q_{k}}$$
,  $k=1,...,M.$  (2.5)

Since  $X_M = Q_M R_M$  we have  $Q_M^T X_M = Q_M^T Q_M R_M = D_M R_M$ , an upper triangular matrix, which gives

$$\mathbf{q}_{\mathbf{k}}^{\mathsf{T}} \mathbf{x}_{\mathbf{j}} = \mathbf{0}$$
 ,  $\mathbf{j} < \mathbf{k}$ 

Finally,  $Q_{M} = X_{M}R_{M}^{-1} = \chi_{M}(\hat{L}_{M}^{-1})^{T}$ , <u>i.e</u> by (2.4),

$$q_k = x_k + \sum_{j=1}^{k-1} \hat{\alpha}_{k-1}(j) x_{k-j}$$
,

which we have shown under the induction hypothesis is  $x_{(k)}$ .

Thus,

$$\hat{a}(k) = -\frac{\sum_{i=1}^{T} \hat{q}_{k}^{T}(j) \times j}{\sum_{i=1}^{T} \hat{q}_{k}^{T}(k)} = -\frac{q_{k}^{T} \left[ \sum_{i=1}^{T} \hat{q}_{k-1}^{T}(j) \times j \right]}{q_{k}^{T} q_{k}}$$

$$= -\frac{q_{k}^{T} \sum_{i=1}^{T} \hat{q}_{k}^{T}(k)}{q_{k}^{T} q_{k}} = \hat{q}_{k}^{T}(k)$$

by (b) and (2.5).

QED.

# Corollary 2.1

The vectors  $y_{(k)}$  and  $x_{(k)}$  are the forward and backward prediction errors one would obtain using a memory (k-1) linear prediction filter assuming  $y(1-k) = \ldots = y(0) = y(T+1) = \ldots = y(T+k) = 0$ , i.e if we denote by  $y_{(k)}^j$  and  $x_{(k)}^j$  the  $j^{th}$  elements of  $y_{(k)}$  and  $x_{(k)}^j$ , then for  $j=1,\ldots,T$ ,

$$y_{(k)}^{j} = -\sum_{\ell=1}^{k-1} \hat{\alpha}_{k-1}(\ell)y(j-\ell)$$

$$x_{(k)}^{j} = -\sum_{k=1}^{k-1} \hat{\alpha}_{k-1}(k)y(j+k)$$

where  $\hat{\alpha}_{k-1}$  is the value of  $a = (a(1),...,a(k-1))^T$  minimizing both

$$S_{F}(a) = \sum_{j=1}^{T} \{y(j) + \sum_{\ell=1}^{k-1} a(\ell)y(j-\ell)\}^{2}$$

and

$$S_{B}(a) = \sum_{j=1}^{T} \{y(j) + \sum_{\ell=1}^{k-1} a(\ell)y(j+\ell)\}^{2}$$
,

assuming y(1-k)=...=y(0)=y(T+1)=...=y(T+k)=0. Further,

$$x_{(k)}^{T}x_{(k)} = s_{B}(\hat{x}_{k-1}) = s_{F}(\hat{x}_{k-1}) = y_{(k)}^{T}y_{(k)}, \qquad (2.6)$$

k=1,...,M+1.

The Toeplitz Gram-Schmidt Yule-Walker algorithm for finding  $\hat{\alpha}_k(j), \hat{\sigma}_k^L, \ 1 \leq j \leq k \leq M, \ \text{consists then of the following steps:}$ 

i) 
$$y_{(1)} = y_M, x_{(1)} = L(y_M), \hat{\sigma}_0^2 = \hat{R}(0)$$
 (2.7)

ii) for k=1,...,M: (2.8)
$$\hat{\alpha}_{k}(k) = -\frac{y(k)^{\times}(k)}{T}$$

$$\frac{\hat{\alpha}_{k}(k)}{\hat{\alpha}_{k}(k)} = \frac{y(k)^{\times}(k)}{T}$$

$$\hat{\sigma}_{k}^{2} = \hat{\sigma}_{k-1}^{2} (1 - \hat{\alpha}_{k}^{2}(k))$$

$$y(k+1) = y(k) + \hat{\alpha}_{k}^{(k)}(k)$$

$$x(k+1) = L(x(k) + \hat{\alpha}_{k}^{(k)}(k))$$

iii) use (2.1), (2.2) to obtain the other  $\hat{\alpha}_k(j)$ 's.

#### 3. NUMERICAL PROPERTIES OF THE ALGORITHM

## 3.1 Arithmetic Operations and Required Storage

To obtain  $\hat{\theta}_{M}$  there are 2(M-1) updates of the form a - cb for (T+M) x l vectors a and b and 2M inner products of (T+M) x l vectors. Counting a multiplication and addition as an operation, there are thus (T+M)(4M-2) operations to obtain  $\hat{\theta}_{M}$ .

To use Levinson's algorithm,  $\hat{R}(0),\ldots,\hat{R}(M)$  must first be calculated, which takes approximately TM operations if inner products are used or 2T  $\log_2$  T operations if the Fast Fourier Transform is used.

In terms of speed then the algorithm of Theorem 2.1 is slightly slower than using Levinson's algorithm. Neither of the algorithms requires significant storage; approximately 2 (T+M) locations for the Toeplitz Gram-Schmidt algorithm versus T+M for Levinson's algorithm when the data vector is stored.

## 3.2 Error Analysis

To determine the numerical accuracy of the proposed algorithm we shall first prove that it is a method of finding the matrix Q and the last column of R in the MGSD of  $[X_M:Y_M]$  using only two vectors of length T+M.

Let A be a full rank (n x p) matrix having columns  $a_1, \ldots, a_p$ . The MGSD algorithm is a numerically stable (Björck (1967)) method of determining the Gram-Schmidt factors  $Q=(q_1,\ldots,q_p)$  and R of A in p stages. At the  $k^{th}$  stage, the  $k^{th}$  column of Q and the  $k^{th}$  row of R are determined. Conceptually, one defines the sequence of matrices  $Q^{(0)}=A$ ,  $Q^{(1)},\ldots,Q^{(p)}=Q$ , with  $Q^{(k)}$  having columns  $q_1^{(k)},\ldots,q_p^{(k)}$ , by:

$$q_{j}^{(k)} = \begin{cases} q_{j}^{(k-1)}, & j \leq k \\ q_{j}^{(k-1)} + R_{kj} q_{k}^{(k)}, & j > k \end{cases}$$

where  $R_{kj}$  is the ratio of inner products

$$R_{kj} = -\frac{(g_{j}^{(k-1)})^{T} g_{k}^{(k)}}{(g_{k}^{(k)})^{T} g_{k}^{(k)}}$$

Then  $q_k = q_k^{(k)}$  and one can actually accumulate  $Q^{(0)}$ ,  $Q^{(1)}$ ,..., $Q^{(p)}$  in a single matrix.

We have

## Theorem 3.1

The vectors  $x_{(k)}$  and  $y_{(k)}$  in Theorem 2.1 are the  $k^{th}$  and  $(M+1)^{st}$  columns of  $Q^{(k)}$  in the MGSD of matrix  $[X_M:Y_M]$ .

## Proof

Parts (b) and (c) of Theorem 2.1 show that for k>1,  $y_{(k)}$  and  $x_{(k)}$  and the ordinary least squares residuals of the regression of  $y_{M}$  on  $x_{1}, \dots, x_{k-1} \text{ and } x_{k} \text{ on } x_{k-1}, \dots, x_{1} \text{ respectively and thus (see Clayton (1971)) they are } y_{M+1}^{(k)} \text{ and } y_{k}^{(k)} \text{ respectively. For } k=1,$ 

$$R_{12} = -\frac{x_2^T x_1}{x_1^T x_1} = -\frac{y_M^T x_1}{x_1^T x_1} = R_{1,M+1} = \hat{\alpha}_1(1).$$

QED

Since the Toeplitz Gram-Schmidt algorithm is a subset of the MGSD we can use the results of Björck (1967). We use the techniques and notation introduced by Wilkinson (1965).

Using equation (5.8) from Björck (1967) and assuming that all inner products are accumulated in double precision, we have

$$||fl(y_{(M)}) - y_{(M)}|| \le 1.9 M^{\frac{1}{2}} (M+1)2^{-t} \sqrt{R(0)}$$
, (3.1)

where  $2^{-t}$  is the machine precision of the computer being used to perform the calculations. This proves that the Toeplitz Gram-Schmidt algorithm gives a numerically stable prewhitening filter. By symmetry, a formula identical to (3.1) would apply for  $x_{(M)}$ .

The roundoff error for  $\hat{\alpha}_k(k)$  can be obtained by letting  $\tilde{\alpha}_k(k)$  be the exact solution (no rounding errors) using the inexact  $fl(x_{(k)})$  and  $fl(y_{(k)})$  so that

$$\begin{split} \left| \operatorname{fl} \left( \hat{\alpha}_{k}(k) \right) - \hat{\alpha}_{k}(k) \right| &\leq \left| \operatorname{fl} \left( \hat{\alpha}_{k}(k) \right) - \tilde{\alpha}_{k}(k) \right| \\ &+ \left| \tilde{\alpha}_{k}(k) - \hat{\alpha}_{k}(k) \right| \end{split}$$

Using a perturbation bound on  $\tilde{\alpha}_{k}(k)$  and equation (4.11) in Björck, one obtains

$$|f|(\hat{\alpha}_{k}(k))|-\tilde{\alpha}_{k}(k)|<(2.01|\hat{\alpha}_{k}(k)|+.01\hat{\sigma}_{k}^{2})2^{-t}$$
,

$$|\tilde{a}_{k}(k) - \hat{a}_{k}(k)| < 8k^{\frac{1}{2}}(k+1) \sqrt{R(0)}$$
 2<sup>-t</sup>,

and hence

$$|f|(\hat{\alpha}_{k}(k)) - \hat{\alpha}_{k}(k)| < (2.01|\hat{\alpha}_{k}(k)| + .01|\hat{\sigma}_{k}^{2} + 8k^{\frac{1}{2}}(k+1)|\sqrt{R(0)}|)2^{-t}|.$$

From this it is concluded that the algorithm is stable.

To illustrate the numerical accuracy of the algorithm, we use the example of a constant y(t) (see Davis and Qualls (1977)). For T=1000 and M=20,  $\hat{\Gamma}_{\text{M}}$  has condition number K( $\hat{\Gamma}_{\text{M}}$ ) = 505. The results of using Levinson's algorithm and the Toeplitz Gram-Schmidt algorithm (multiplied by 2<sup>t</sup> to make them reasonably machine independent) are presented in Table 3.1.

	Table 3.1	
	$  f (\hat{a}_{M})-\hat{a}_{M}  2^{t}$	K
Levinson	1168	505
Toeplitz Gram-Schmidt	. 41	22.5

#### 4. AUTOREGRESSION DIAGNOSTICS AND ROBUST AUTOREGRESSION

In this section we consider the application of methods in ordinary regression analysis for robustness and detecting influential observations in the autoregression situation. We will show that the Toeplitz Gram-Schmidt algorithm provides a unified framework for carrying out this analysis.

## 4.1 Detecting Influential Observations

In the usual full rank, n-observation, p-parameter, linear model  $y = X\beta + \varepsilon$ , the diagonal elements of the hat matrix  $H = X(X^TX)^{-1}X^T$  are often used (Hoaglin and Welsch (1978)) to determine if undue weight is given any particular observation in finding the vector of predicted values  $\hat{y} = X\hat{\beta} = X(X^TX)^{-1}X^Ty = Hy$ .

If X = QR where  $Q^TQ = D = Diag(d_1, \dots, d_p)$ , we have  $H = QR[R^{-1}D^{-1}R^{-T}]R^TQ^T = QD^{-1}Q^T$ . Thus

$$H_{jj} = \sum_{k=1}^{p} Q_{jk/d_k}^2$$
, j=1,...,n

In the autoregression case then these hat diagonals can be easily accumulated in the Toeplitz Gram-Schmidt Algorithm. If we define  $d_{(k)} = \sum_{k=1}^{T} X_{(k)} X_{(k)}, \quad k=1,\dots,M+1, \text{ and}$ 

$$h_{(1)}^{j} = (x_{(1)}^{j})^{2}/d_{(1)}$$
,  $j=1,...,T+M$ ,

$$h_{(k)}^{j} = h_{(k-1)}^{j} + (x_{(k)}^{j})^{2}/d_{(k)}$$
,  $j=1,...,T+M,k=2,...,M+1$ ,

then  $h_{(k)} = (h_{(k)}, \dots, h_{(k)}^{T+M})^T$  are the hat diagonals of the regression of  $y_M$  on  $-x_1, \dots, -x_{k-1}$ . As such they can be used both for detecting influential observed y's and for measuring end effects. Note that calculation of the  $h_{(k)}$  can be done easily by just calculting  $d_{(k)}$  and  $h_{(1)}$  in (2.7) and  $d_{(k)}, h_{(k)}$  in (2.8).

#### 4.2 Alternative Estimators

A wide variety of possibly robust estimators of autoregressive parameters are suggested by the results of Theorem 2.1 and Corollary 2.1, i.e the Yule-Walker  $\hat{\alpha}_k(k)$  is the regression coefficient of the regression through the origin of the forward prediction errors  $\underline{y}_{(k)}$  on the backward prediction errors  $\underline{x}_{(k)}$ . Thus any of the M-estimators (Huber (1973) could be used rather than ordinary least squares to obtain estimators  $\tilde{\alpha}_1(1),\ldots,\tilde{\alpha}_M(M)$ . Then these would be used instead of  $\hat{\alpha}_1(1),\ldots,\hat{\alpha}_M(M)$  in (2.1)-(2.3), thus giving what could be called  $\hat{x}_2$ -M autoregressive parameter estimators,  $\hat{x}_2$  for the use of Levinson's algorithm and M for the use of M-estimators. We note that such M-estimators probably need not lead to  $\tilde{\alpha}_k(k)$ 's that are less than one in absolute value (A necessary and sufficient condition for stability of resulting prediction filters) but this may actually be a benefit in that it would indicate that the original time series is not stationary.

Now since  $x_{(k)}^T = y_{(k)}^T = 0$  if  $y_{M}^T = 0$  we have also that  $\hat{\alpha}_k(k)$  is the negative of the ordinary sample correlation coefficient of the forward and backward prediction errors for order (k-1). Thus again one could use robust methods of estimating correlation coefficients (Huber (1977)) to obtain other estimators  $\tilde{\alpha}_k(k)$ .

## 4.3 Relationship of the Toeplitz Gram-Schmidt and Burg Algorithms

An important method of autoregressive parameter estimation is the Burg (1968) algorithm. With the framework provided by the Toeplitz Gram-Schmidt algorithm it is easy to describe what the Burg algorithm does. It uses the same updating equations but uses

$$\tilde{\alpha}_{k}(k) = -\sum_{j=1}^{T-k} y_{(k)}^{j+k} x_{(k)}^{j+k}$$

$$\frac{1}{2} \left[ \sum_{j=1}^{T-k} (x_{(k)}^{j+k})^{2} + \sum_{j=1}^{T-k} (y_{(k)}^{j+k})^{2} \right]$$

instead of  $\hat{\alpha}_k(k)$ . By (2.6) note that  $\tilde{\alpha}_k(k)$  is identical to  $\hat{\alpha}_k(k)$  except that the summations range from 1 to T-k instead of from 1 to T+M. Note also that  $\tilde{\alpha}_k(k)$  is the value of a minimizing

$$\sum_{j=1}^{T-k} \{ (y_{(k)}^{j+k} + a x_{(k)}^{j+k})^2 + (x_{(k)}^{j+k} + a y_{(k)}^{j+k})^2 \},$$

i.e the sum of squares of error in regressing the "observed" forward prediction errors on the "observed" backward prediction errors and vice versa, i.e the Burg algorithm never uses a youtside the range of observed values. We note further that Stuart (1955) has shown that if  $(x_1,y_1),\ldots,(x_n,y_n)$  is a random sample from a bivariate normal distribution with zero means and equal but unknown variances, then

$$r' = \frac{\sum_{i=1}^{n} x_{i} y_{i}}{\frac{1}{2} \left[ \sum_{i=1}^{n} x_{i}^{2} + \sum_{i=1}^{n} y_{i}^{2} \right]}$$

is an asymptotically efficient estimator of the correlation coefficient of x and y. Thus the Burg estimator of  $\alpha_k(k)$  can also be interpreted as the negative of a correlation coefficient estimator.

Thus if k is large relative to T and R(0) is small the Burg algorithm may be inferior to Yule-Walker estimation, while if R(0) is large it will probably be superior because of end effects in Yule-Walker estimation. Finally we note that the  $\tilde{\alpha}_k(k)$  are guaranteed to be less than one in absolute value.

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